

OM nucleic - nucleic search, using sw model

Run on: June 2, 2003, 00:51:28 ; Search time 4674.33 Seconds
(without alignments)
16997.243 Million cell updates/sec

Title: US-09-596-194-59
Perfect score: 2730
Sequence: 1 gtcgacccacgcgtccgtcc.....aaaaaaaaaagggcgccgc 2730

Scoring table: OLIGO_NUC
Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 400

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:*
1: gb_ba:*
2: gb_htg:*
3: gb_in:*
4: gb_om:*
5: gb_ov:*
6: gb_pat:*
7: gb_ph:*
8: gb_pl:*
9: gb_pr:*
10: gb_ro:*
11: gb_sts:*
12: gb_sy:*
13: gb_un:*
14: gb_vi:*
15: em_ba:*
16: em_fun:*
17: em_hum:*
18: em_in:*
19: em_mu:*
20: em_om:*
21: em_or:*
22: em_ov:*
23: em_pat:*
24: em_ph:*
25: em_pl:*
26: em_ro:*
27: em_sts:*

search repeated in
GenSeq
US Pats
US P&PUBS
EST

GenCore version 5.1.4_p5_4578
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Run on: June 2, 2003, 00:51:28 ; Search time 3446.67 Seconds
(without alignments)
16997.243 Million cell updates/sec

Title: US-09-596-194-60
Perfect score: 2013
Sequence: 1 atggcccagctgttcctgcc.....ccctgcagctctgggtcacc 2013

Scoring table: OLIGO_NUC
Gapop 60.0 , Gapext 60.0

Searched: 2054640 seqs, 14551402878 residues

Word size : 400

Total number of hits satisfying chosen parameters: 9

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Listing first 45 summaries

Database : GenEmbl:*
1: gb_ba:*
2: gb_htg:*
3: gb_in:*
4: gb_om:*
5: gb_ov:*
6: gb_pat:*
7: gb_ph:*
8: gb_pl:*
9: gb_pr:*
10: gb_ro:*
11: gb_sts:*
12: gb_sy:*
13: gb_un:*
14: gb_vi:*
15: em_ba:*
16: em_fun:*
17: em_hum:*
18: em_in:*
19: em_mu:*
20: em_om:*
21: em_or:*
22: em_ov:*
23: em_pat:*
24: em_ph:*
25: em_pl:*
26: em_ro:*
27: em_sts:*

Search repeated in
GenSeq
US Pats
US PGPubs
EST